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OM protein nucleic search, using frame_plus_p2n model

Run on: January 16, 2003, 16:51:22 : Search time 212.829 Seconds
(without alignments)
137.557 Million cell updates/sec

Title: US-09-856-070-19

Perfect score: 65

Sequence: 1 KKEIMLRIDYEE 13

Scoring table: ELASOM62

Xgapop 10.0, Xgapext 0.5

Ygapop 10.0, Ygapext 0.5

Zgapop 6.0, Zgapext 7.0

Delgap 6.0, Delgapext 7.0

Searched: 2185236 seqs, 113000160 residues

Total number of hits satisfying chosen parameters: 4370478

Minimum DB seq length: 0

Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Command line parameters:

-MODEL=frame_p2n model -DEFV=xlfp
-Q=/cgc2.1/seqpt0_spool/US09856070/cunat_14012003_158433_1611/app_query.fasta_1_1592
-PR=N_Genescs_1_161002 -GRMI=fastaq -SUFIX=ref -MINMATCH=0.1 -LOG=1-0
-LOCAL=LOCAL -OUTFMT=ptn -NRM=ext -HEAPSIZE=500 -MINLEN=100 -THRESH=0 -ALIGN=15
-USEP=US09856070/seqpt0_1_14444_sufat_14012003_158433_1611 -NPROG 6 -TOPIC 4
-NO_XLIFY -NO_MMAP -LAPGENEPE -NEW_SCORES 0 -WAIT -LONGEST -REV -TIMEOUT 120
-WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -ZGAPOP=6 -ZGAPEXT=7
-YGAPOP=10 -YGAPEXT=0.5 -DELAP=6 -DELAPEXT=7

Database: N_Genescs_101002:*

1: /SID52/qcdata/qcdata/geneseq/geneseq-emb1/NA1980.DAT:*

2: /SID52/qcdata/qcdata/geneseq/geneseq-emb1/NA1981.DAT:*

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19: /SID52/qcdata/qcdata/geneseq/geneseq-emb1/NA1998.DAT:*

20: /SID52/qcdata/qcdata/geneseq/geneseq-emb1/NA1999.DAT:*

21: /SID52/qcdata/qcdata/geneseq/geneseq-emb1/NA2000.DAT:*

22: /SID52/qcdata/qcdata/geneseq/geneseq-emb1/NA2001.DAT:*

23: /SID52/qcdata/qcdata/geneseq/geneseq-emb1/NA2001B.DAT:*

24: /SID52/qcdata/qcdata/geneseq/geneseq-emb1/NA2002.DAT:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	65	100.0	2595	22	Human colon cancer
2	65	100.0	2930	24	Human osteoblast d
3	65	100.0	2930	24	Human lung cancer
4	65	100.0	3044	24	Human osteoblast d
5	65	100.0	3044	24	Human cDNA differe
6	65	100.0	3044	24	Cene #1721 used to
7	65	100.0	3047	24	Human ovarian tumor
8	65	100.0	4072	24	Human osteoblast d
9	65	100.0	3115	21	Human colon cancer
10	65	100.0	11445	22	Human immune/haema
11	41	63.1	1447	23	DNA encoding novel
12	41	63.1	6694	24	Human osteoblast d
13	40	61.5	224	22	Human foetal liver
14	40	61.5	224	22	Human brain expres
15	40	61.5	520	22	Human foetal liver
16	40	61.5	520	22	Human brain expres
17	40	61.5	1815	23	DNA encoding novel
18	40	61.5	2528	21	Human secreted pro
19	40	61.5	2701	22	Human cDNA sequen
20	40	61.5	2901	24	Human ovarian anti
21	40	61.5	2979	23	Human prostate exp
22	40	61.5	4226	23	DNA encoding novel
23	40	61.5	4558	22	Human cervical can
24	40	61.5	149671	24	Human cDNA differe
25	39	60.0	205	22	Human foetal liver
26	39	60.0	205	22	Human brain expres
27	39	60.0	205	22	Human bone marrow
28	39	60.0	205	22	Probe #20109 used
29	39	60.0	205	22	Human genome deriv
30	39	60.0	205	22	Human secreted pro
31	39	60.0	553	24	Human colon tumor
32	39	60.0	592	22	DNA encoding G pro
33	39	60.0	1256	21	Human low adenosin
34	39	60.0	1256	21	Human adenosine re
35	39	60.0	1400	13	Encodes a HeLa cel
36	39	60.0	1400	13	DNA encoding a gly
37	39	60.0	1867	22	Human colon cancer
38	39	60.0	1985	23	DNA encoding novel
39	39	60.0	2134	18	Human myeloid cell
40	39	60.0	2142	23	DNA encoding novel

ALIGNMENTS

RESULT 1

AAH33385

12 AAH33385 standard. LNA. 2595 BP.

XX AAH33385;

XX 03 Sep 2001 (first entry)

XX Human colon cancer antigen encoding cDNA SEQ ID NO:441.

XX Homo sapiens, cell carcinoma, detection;

XX colorectal carcinoma; SS.

XX Homo sapiens.

XX W250122920 A2.

XX 05-APR-2001.

XX 28-SEP-2000; 2000W0-US26424.
 XX 29-SEP-1999; 9909-0157147
 XX 03-NOV-1999; 9909-0163280.
 XX (HUMAN) HUMAN GENOME SCI INC.
 XX Ruben SM, Barash SC, Birse CE, Rosen CA;
 XX WPI; 2001-235357/24.
 XX P-PSDB; AAG73954.
 XX Nucleic acids encoding 4277 human colon cancer-associated polypeptides,
 XX useful for preventing, diagnosing and/or treating colorectal cancers -
 XX Claim 1; Page 2539-2540; 9803pp; English.
 XX AAH32943 to AAH37195 and AAH373514 to AAH37788 represent human colon
 XX cancer-associated nucleic acid molecules (N) and proteins (P), where
 XX the proteins are collectively known as colon cancer antigens. The colon
 XX cancer antigens have cytostatic activity and can be used in gene
 XX therapy and vaccine production. N and P may be used in the prevention,
 XX diagnosis and treatment of diseases associated with inappropriate P
 XX expression. For example, N and P may be used to treat disorders
 XX associated with decreased expression by rectifying mutations or deletions
 XX in a patient's genome that affect the activity of P by expressing
 XX inactive proteins or to supplement the patients own production of P.
 XX Additionally, N may be used to produce the colon cancer-associated ps,
 XX by inserting the nucleic acids into a host cell and culturing the cell
 XX to express the proteins. N and P can be used in the prevention, diagnosis
 XX and treatment of colorectal carcinomas and cancers. AAH37196 to AAH37234
 XX and AAH37789 represent sequences used in the exemplification of the
 XX present invention.
 XX N.B. Pages 666 to 682 and page 7053 of the sequence listing were
 XX missing at time of publication, meaning no sequences are present for
 XX SEQ ID NO:1027 to 1052, 7921 and 7922.
 XX SQ Sequence 2595 BP; 742 A; 562 C; 714 G; 567 T; 10 other;
 Alignment Scores:
 Pred. No.: 0.00473 Length: 2595
 Score: 65.00 Matches: 13
 Percent Similarity: 100.00% Conservative: 0
 Best Local Similarity: 100.00% Mismatches: 0
 Query Match: 100.00% Indels: 0
 DB: 22 Gaps: 0
 US-09-856-070-19 (1-13) x AAH33385 (1-2595)
 QY 1 LysGluGluLeuMetLeuArqLeuGlnAspTyrGluGlu 13
 DB 658 AAGGAGAGTGTGCTGCTGCGGCTGCAGGACTATGAGGAG 696
 RESULT 2
 AAH88181
 ID AAH88181 standard; cDNA: 2930 BP.
 XX AAH88181;
 AC AAH88181;
 XX 18-SEP-2002 (first entry)
 DE Human osteoblast differentiation related cDNA SEQ ID NO 88.
 XX Human; osteoblast; stem cell differentiation; bone tissue deposition;
 XX osteoporosis; osteopathic; ss.
 XX Homo sapiens.
 XX W0200250401-A2
 XX 27-JUN-2002.
 XX

PF 18-DEC-2001; 2001W0-US49476.
 XX 18-DEC-2000; 2000TS-255882P.
 XX 24-APR-2001; 2001TS-285691P.
 XX (GENE-) GENE LOGIC INC.
 XX PA (PROC) PROCTER & GAMBLE CO.
 XX Ji D, Axelrod LW, Cook JS, Jaiswal N, Einstein P, Houghton A;
 XX Mertz L;
 XX WPI; 2002-557663/59.
 XX Use of genes and their expression profiles associated with osteoblast
 XX differentiation for screening modulators bone formation, for diagnosing
 XX or treating e.g. osteoporosis, or as markers for the differentiation
 XX process -
 XX Claim 1, SEQ ID NO 88, 78pp; Sequence listing: English.
 XX The invention relates to genes and their expression profiles are used
 XX for:
 XX (a) screening modulators of precursor stem cell differentiation into
 XX osteoblasts, or bone tissue deposition;
 XX (b) diagnosing abnormal deposition of bone tissue, abnormal rate of
 XX osteoblast formation or osteoporosis; or
 XX (c) treating or monitoring treatment of the conditions cited in (b), or
 XX monitoring the progression of bone tissue deposition.
 XX Specific conditions include postmenopausal osteoporosis, glucocorticoid
 XX osteoporosis or male osteoporosis, osteopenia, osteodystrophy,
 XX drug-induced abnormalities in bone formation or bone loss, conditions
 XX that involve altered bone metabolism (e.g. idiopathic juvenile
 XX osteoporosis), skeletal disease linked to breast cancer, mastocytosis,
 XX Fanconi syndrome or fibrous dysplasia. The present sequence is that of an
 XX osteoblast differentiation associated cDNA marker of the invention.
 XX Note: The sequence data for this patent did not form part of the printed
 XX specification, but was obtained in electronic format directly from WIPO
 XX at ftp.wipo.int/pub/published_pat_sequences.
 XX SQ Sequence 2930 BP; 793 A; 658 C; 821 G; 658 T; 0 other;
 Alignment Scores:
 Pred. No.: 0.00545 Length: 2930
 Score: 65.00 Matches: 13
 Percent Similarity: 100.00% Conservative: 0
 Best Local Similarity: 100.00% Mismatches: 0
 Query Match: 100.00% Indels: 0
 DB: 24 Gaps: 0
 US-09-856-070-19 (1-13) x AAH88181 (1-2930)
 QY 1 LysGluGluLeuMetLeuArqLeuGlnAspTyrGluGlu 13
 DB 1106 AAGTACGACTTTCATGCTGCTGCGGCTGCAGGACTATGAGGAG 1144
 RESULT 3
 AAH70285
 ID AAH70285 standard; cDNA: 2930 BP.
 XX AAH70285;
 AC AAH70285;
 XX 15-JUL-2002 (first entry)
 DE Human lung cancer associated full length cDNA DMSM-51.
 XX Human; ss, gene; lung cancer; cytostatic; tumour; vaccine.
 XX Homo sapiens.
 XX W0200224057 A2
 XX 28-MAR-2002.
 XX

PF 20-SEP-2001; 2001WO 0542232.
 XX
 PP 22-SEP-2000; 2000PS-234887P
 PP 10-OCT-2000; 2000US-249449P.
 PP 20-JUN-2001; 2001US-301928P.
 XX
 PA (COR1-) COMEXA COMP.
 PA
 PI Benson DP, Mohanath P, Lodes MT;
 XX
 XX WPI: 2002 372001/40
 XX
 XX New tumor lung proteins and nucleic acids encoding the proteins, useful
 PI as vaccines and for treating, preventing, diagnosing or monitoring lung
 PT cancer
 PT
 XX
 PS Claim 1: Page 159-160, 189pp; English.
 XX
 CC The invention relates to an isolated polynucleotide comprising a sequence
 CC selected from 183 human DNA sequences (appearing as AHK70130-AHK70312),
 CC or their fragments, homologues, variants or complements and their encoded
 CC polypeptides. Also included are an expression vector comprising the
 CC polynucleotide operably linked to an expression control sequence; a host
 CC cell transformed or transfected with an expression vector; an isolated
 CC antibody, or its antigen-binding fragment that specifically binds to the
 CC polypeptide; a method for detecting the presence of a cancer in a
 CC patient; a fusion protein comprising at least the polypeptide; an
 CC oligonucleotide that hybridises to the polynucleotide under moderately
 CC stringent conditions; a method for stimulating and/or expanding T cells
 CC specific for a tumour protein; an isolated T cell population comprising T
 CC cells prepared from the method of above; a composition comprising a first
 CC component consisting of carriers and immunostimulants, and a second
 CC component selected from the polynucleotides, proteins, antibodies, fusion
 CC proteins, T cell populations and antigen presenting cells expressing the
 CC polypeptide; methods for stimulating an immune response or treating
 CC cancer in a patient by administering the composition and diagnostic kits
 CC comprising at least one of the oligonucleotide of, or an antibody and a
 CC detection reagent consisting of a reporter group. The polypeptides and
 CC polynucleotides are useful as vaccines for the treatment or prevention of
 CC lung cancer, and for diagnosis and monitoring of such cancer. The
 CC polynucleotide, polypeptide and antigen presenting cells can be
 CC used to stimulate or expand T cells specific for a tumorous protein.
 CC The polynucleotides may be used as probes or primers for nucleic acid
 CC hybridisation, and in the preparation of ribozyme molecules for
 CC inhibiting expression of tumour polypeptides and proteins in tumour
 CC cells. The present sequence is one of the 183 lung cancer associated
 CC polynucleotides.
 XX
 SQ Sequence 2930 BP, 793 A; 658 C; 821 G; 658 T; 0 other;

Alignment Scores:
 Pred. No.: 0.00545 Length: 2940
 Score: 65.00 Matches: 13
 Percent Similarity: 100.00% Conservative: 0
 Best Local Similarity: 100.00% Mismatches: 0
 Query Match: 100.00% Indels: 0
 DB: 24 Gaps: 0

US-09-856-070-19 (1-13) x AHK70285 (1-2940)

OY 1 LysGluGluGluMetLeuArgGluGlnAspTyrGluGlu 13
 Db 1106 AAGGAGGATTAATGTTGAGGATGAGGATATATAGGAG 1144
 |||||||

RESULT 4
 AHQ88180
 ID ABQ88180 standard; cDNA: 3044 BP.
 XX
 AC ABQ88180;
 XX
 XX 18-SEP-2002 (first entry)

DT Human osteoblast differentiation related cDNA SEQ ID NO 87.
 DE

XX Human, osteoblast, stem cell differentiation; bone tissue deposition;
 KW osteoporosis; osteopathic; ss.
 XX
 OS Homo sapiens.
 XX
 PN WO200258301 A2.
 XX
 PD 27-JUN-2002.
 XX
 PF 18 DEC-2001, 2001WO-0548276.
 XX
 XX 18-DEC-2000; 2000US 255822P.
 PP 24-APR-2001; 2001US 285691P.
 XX
 PA (GENE-) GENE LOGIC INC.
 PA (PRO-) PRACER & GAMBLE CO.
 XX
 PI Ji D, Axelrod DW, Cook JS, Jaiswal N, Einstein K, Houghton A;
 PI Mertz L;
 XX
 DR WPI: 2002-557663/59.
 XX
 XX Use of genes and their expression profiles associated with osteoblast
 PT differentiation for screening modulators bone formation, for diagnosing
 PT or treating e.g. osteoporosis, or as markers for the differentiation
 XX process
 XX
 PS claim 1; SEQ ID NO 87; 78pp + Sequence Listing; English.
 XX
 CC the invention relates to genes and their expression profiles are used
 CC for:
 CC (a) screening modulators of precursor stem cell differentiation into
 CC osteoblasts, or bone tissue deposition;
 CC (b) diagnosing abnormal deposition of bone tissue, abnormal rate of
 CC osteoblast formation or osteoporosis; or
 CC (c) treating or monitoring treatment of the conditions cited in (b), or
 CC monitoring the progression of bone tissue deposition.
 CC Specific conditions include postmenopausal osteoporosis, glucocorticoid
 CC osteoporosis or male osteoporosis, osteopenia, osteodysraphy,
 CC drug-induced abnormalities in bone formation or bone loss, conditions
 CC that involve altered bone metabolism (e.g. idiopathic juvenile
 CC osteoporosis), skeletal disease linked to breast cancer, mastocytosis,
 CC Fanconi syndrome or fibrous dysplasia. The present sequence is that of an
 CC osteoblast differentiation associated cDNA marker of the invention.
 CC Note: The sequence data for this patent did not form part of the printed
 CC specification, but was obtained in electronic format directly from WIPO
 CC at ftp.wipo.int/pub/published_pct_sequences.
 XX
 SQ Sequence 3044 BP; 826 A; 687 C; 855 G; 675 T; 1 other;

Alignment Scores:
 Pred. No.: 0.0057 Length: 3044
 Score: 65.00 Matches: 13
 Percent Similarity: 100.00% Conservative: 0
 Best Local Similarity: 100.00% Mismatches: 0
 Query Match: 100.00% Indels: 0
 DB: 24 Gaps: 0

US-09-856-070-19 (1-13) x AHQ88180 (1-3044)

OY 1 LysGluGluGluMetLeuArgGluGlnAspTyrGluGlu 13
 Db 1147 AAGGAGGATTAATGTTGAGGATGAGGATATATAGGAG 1185
 |||||||

RESULT 5
 ABK84552
 ID ABK84552 standard; cDNA: 3044 BP.
 XX
 AC ABK84552;
 XX
 XX 14-AUG-2002 (first entry)

DT Human osteoblast differentiation related cDNA SEQ ID NO 87.
 DE

DE Human cDNA differentially expressed in granulocytic cells #1123.
 XX Human, ss: granulocytic cells; DNA chip; bacterial infection;
 KW viral infection; parasitic infection; protozoal infection;
 KW fungal infection; sterile inflammatory disease; psoriasis;
 KW rheumatoid arthritis; glomerulonephritis; asthma; thrombosis;
 KW cardiac reperfusion injury; renal reperfusion injury; AIDS;
 KW adult respiratory distress syndrome; inflammatory bowel disease;
 KW Crohn's disease; ulcerative colitis; periodontal disease;
 KW granulocyte activation; chronic inflammation; allergy.
 XX Homo sapiens.
 OS
 XX
 XX W0200228999-A2.
 XX
 XX 11-APR-2002.
 XX
 XX 03-OCT-2001; 2001W0 0330921.
 XX
 XX 03-OCT-2000; 2000US-247149P.
 XX
 XX (GENE-) GENE LOGIC INC.
 XX
 XX Razzer Barclay Y, Weissman SM, Yamaga S, Vockley J;
 XX W01: 2002-43532R/46
 XX
 XX Detecting granulocyte activation by detecting differential expression
 XX of genes associated with granulocyte activation, which serves as
 XX diagnostic markers that is useful for monitoring disease states and
 XX drug toxicity -
 XX
 XX Claim 1: SEQ ID No 1123; 114pp; English.
 XX
 XX The invention relates to detecting (M1) granulocyte (G) activation
 XX (GCA), by detecting the level of expression of gene(s) (Gs) identified by
 XX DNA chip analysis as given in the specification, and comparing
 XX the expression level to an expression level in an unactivated
 XX Gc, where differential expression of Gs is indicative of GCA.
 XX Also included are modulating (M2) Gc by contacting Gc with an agent
 XX that alters the expression of at least one gene in Gs; (2) screening (M3)
 XX for an agent capable of modulating GCA or an inflammation (especially
 XX chronic) in a tissue, an allergic response in a subject, exposure of a
 XX subject to a pathogen or sterile inflammatory disease using the
 XX gene expression profile; (3) detecting (M4) an inflammation (especially
 XX chronic) in a tissue, an allergic response in a subject, exposure of a
 XX subject to a pathogen or sterile inflammatory disease, by detecting the
 XX level of expression in a sample of the tissue of gene(s) from Gs, where
 XX the level of expression of the gene is indicative of inflammation;
 XX (4) treating (M5) an inflammation (especially chronic) or in a tissue,
 XX an allergic response in a subject, exposure of a subject to a pathogen
 XX or sterile inflammatory disease, by contacting a tissue having
 XX inflammation with an agent that modulates the expression of gene(s)
 XX from Gs in the tissue. M1 is useful for detecting GCA; M2 is useful for
 XX modulating Gc; M3 is useful for screening an agent capable of modulating
 XX GCA preferably in an inflammation in a tissue; M4 is useful for
 XX detecting an inflammation (especially chronic) in a tissue, an allergic
 XX response in a subject, exposure of a subject to a pathogen or sterile
 XX inflammatory disease (e.g. psoriasis, rheumatoid arthritis,
 XX glomerulonephritis, asthma, thrombosis, cardiac reperfusion injury, renal
 XX reperfusion injury, AIDS, adult respiratory distress syndrome,
 XX inflammatory bowel disease, Crohn's disease, ulcerative colitis,
 XX periodontal disease, also bacterial infection, viral infection,
 XX parasitic infection, protozoal infection, fungal infection and M5 is
 XX useful for treating one of the above conditions. The present
 XX sequence represents a gene differentially expressed in granulocytes.
 XX Note: the sequence data for this patent did not form part
 XX of the printed specification, but was obtained in electronic
 XX format directly from WIPO at
 XX http://wipo.int/pub/published_pct_sequences.
 XX
 XX Sequence 3044 BP; 826 A; 687 C; 855 G; 675 T; 1 other;
 XX

Alignment Scores:
 Pred. No.: 0.0057 Length: 3044
 Score: 65.00 Matches: 13
 Percent Similarity: 100.00% Conservative: 0
 Best Local Similarity: 100.00% Mismatches: 0
 Query Match: 100.00% Indels: 0
 DB: 24 Caps: 0
 US-09-856-070-19 (1-13) x ARK#4562 (1-3044)
 QY 1 LysGluGluLeuMetLeuArgLeuGlnAspTyrGluGlu 13
 ID 1147 AAGGAGGAGTTCATGCTGGCGGTGGCAGGACATACAGGAG 1185
 RESULT 6
 ABN97223
 ID ABN97223 standard: DNA; 3044 BP.
 XX
 XX AC ABN97223;
 XX
 XX DT 13-AUG-2002 (first entry)
 XX
 XX DE Gene #3721 used to diagnose liver cancer.
 XX
 XX KW Gene, liver cancer, ds, hepatocellular carcinoma; hepatotropic;
 KW metastatic liver tumor; cytostatic, expression profile; disease state;
 KW disease progression; drug toxicity; drug efficacy; drug metabolism.
 XX
 XX OS Homo sapiens.
 XX
 XX XX W0200229103-A2.
 XX
 XX PD 11-APR-2002.
 XX
 XX PF 02-OCT-2001; 2001W0-US30589.
 XX
 XX PR 02-OCT-2000; 2000US-237054P.
 XX
 XX PA (GENE-) GENE LOGIC INC.
 XX
 XX PI Horne D, Alvares C, Peres-Da-Silva S, Vockley JG;
 XX W01: 2002 426119/45.
 XX
 XX Diagnosing and detecting the progression of liver cancer,
 XX hepatocellular carcinoma or metastatic liver tumor in a patient,
 XX involves detecting the level of expression of two or more genes in a
 XX liver tissue sample -
 XX
 XX Claim 1: SEQ ID NO 3721; 298pp; English.
 XX
 XX The invention relates to a novel method for diagnosing and detecting the
 XX progression of liver cancer, hepatocellular carcinoma or metastatic liver
 XX tumor in a patient, and differentiating metastatic liver cancer from
 XX hepatocellular carcinoma in a patient, involving detecting the level of
 XX expression of two or more genes represented in ARN93503-ARN97455 in a
 XX tissue sample. The method of the invention has hepatotropic, and
 XX cytostatic activity. The method is useful for diagnosing and detecting
 XX the progression of liver cancer, hepatocellular carcinoma and metastatic
 XX liver carcinoma in a patient. The method is useful for identifying
 XX expression profiles which serve as useful diagnostic markers as well as
 XX markers that can be used to monitor disease states, disease progression,
 XX drug toxicity, drug efficacy and drug metabolism.
 XX Note: The sequence data for this patent did not form part of the printed
 XX specification, but was obtained in electronic format directly from WIPO
 XX at http://wipo.int/pub/published_pct_sequences.
 XX
 XX Sequence 3044 BP; 826 A; 687 C; 855 G; 675 T; 1 other;
 XX
 XX Alignment Scores:
 Pred. No.: 0.0057 Length: 3044
 Score: 65.00 Matches: 13
 Percent Similarity: 100.00% Conservative: 0

Best Local Similarity: 100.00%
Query Match: 100.00%
Mismatches: 0
Indels: 0
Gaps: 0

US-09-856-070-19 (1-13) x ARK97223 (1-3044)

QY 1 LysGluGluLeuMetLeuArgLeuGlnAspTyrGluGlu 13
|||||
DB 1147 AAGGAGAGATGATGCTGGCGCTGACGAGACTATGAGGAG 1185

RESULT 7

ABK09792
ID ARK09792 standard; cDNA: 3047 BP

XX AC ARK09792;

XX DT 14-MAR-2002 (first entry)

XX DE Human ovarian tumour protein encoding cDNA #325

XX KW Human; ovarian tumour protein, cancer, cytostatic, immunostimulant, ss,
KW gene therapy; CD4+ T cell, CD8+ T cell, PCR primer.

XX OS Homo sapiens.

XX PN W0200190154-A2

XX PD 29-NOV-2001.

XX PF 23-MAY-2001; 2001WO-051680e

XX PP 24-MAY-2000; 2000US-207107p

XX PP 13-JUN-2000; 2000US-211457p

XX PP 21-JUN-2000; 2000US-211474p

XX PP 03-AUG-2000; 2000US-232288p

XX PP 01-MAR-2001; 2001US-027290p

XX PA (CORI-) CORIXA CORP.

XX PI Xu J, Mitcham JL, Harlocker SL, Dillon DC, Seerist R, Lodes MJ;

XX PI Aude PA, Fling SP, Mannion J, Benson DR, Carter D;
XX DR WPI: 2002-097641/13

XX PS Claim 1; Page 269 270; 285pp; English.

XX CC The invention relates to an isolated polynucleotide encoding a
XX CC polypeptide comprising a portion of an ovarian tumour protein. The
XX CC sequences of the invention are useful for stimulating an immune response
XX CC and for treating ovarian cancer in a patient. An antigen presenting cell
XX CC that expresses the sequences is useful for treating ovarian cancer by
XX CC incubating CD4+ and/or CD8+ T cells isolated from a patient. The T cells
XX CC can then be proliferated and administered to the patient to inhibit the
XX CC development of cancer. The DNA sequences are useful as probes or primers
XX CC for nucleic acid hybridisation, to direct expression of a polypeptide in
XX CC appropriate host cells. Detecting the presence of a cancer in a patient
XX CC involves obtaining a biological sample from the patient, contacting the
XX CC biological sample with an agent that binds to the protein, detecting the
XX CC amount of protein that binds to the agent, comparing the amount of
XX CC protein to a predetermined cut-off value and determining the presence of
XX CC cancer. Sequences ARK09464-ARK09802 represent PCR primers and cDNA
XX CC molecules encoding ovarian tumour proteins of the invention

XX SQ Sequence 3047 BP; 828 A; 687 C; 856 G; 675 T; 1 other;

Alignment Scores:
Pred. No.: 0.0057 Length: 3047
Score: 65.00 Matches: 13
Percent Similarity: 100.00% Conservative: 0

Best Local Similarity: 100.00%
Query Match: 100.00%
Mismatches: 0
Indels: 0
Gaps: 0

US-09-856-070-19 (1-13) x ARK09792 (1-3047)

QY 1 LysGluGluLeuMetLeuArgLeuGlnAspTyrGluGlu 13
|||||
DB 1147 AAGGAGAGATGATGCTGGCGCTGACGAGACTATGAGGAG 1185

RESULT 8

ABK08182
ID ARK08182 standard; cDNA: 3072 BP

XX AC ARK08182;

XX DT 18-SEP-2002 (first entry)

XX DE Human osteoblast differentiation related cDNA SEQ ID NO 89.

XX KW Human; osteoblast, stem cell differentiation, bone tissue deposition;
KW osteoporosis, osteopathic; ss.

XX OS Homo sapiens.

XX PN W0200250301-A2.

XX PD 27-JUN-2002.

XX PF 18-DEC-2001; 2001WO-0548276.

XX PP 18-DEC-2000; 2000US-252822P.

XX PP 24-APR-2001; 2001US-285691P.

XX PA (GENE-) GENE LOGIC INC.

XX PA (PROM) PROMETTER & GAMBLE CO.

XX PI Ji D, Axelrod DW, Cook JS, Jaiswal N, Einstein R, Houghlon A;

XX PI Mertz L;

XX DR WPI: 2002-557663/59.
XX CC The invention relates to genes and their expression profiles are used
XX CC for:
XX CC (a) screening modulators of precursor stem cell differentiation into
XX CC osteoblasts, or bone tissue deposition;
XX CC (b) diagnosing abnormal deposition of bone tissue, abnormal rate of
XX CC osteoblast formation or osteoporosis; or
XX CC (c) treating or monitoring treatment of the conditions cited in (b), or
XX CC monitoring the progression of bone tissue deposition.
XX CC Specific conditions include postmenopausal osteoporosis, glucocorticoid
XX CC osteoporosis or male osteoporosis, osteopenia, osteodystrophy,
XX CC drug-induced abnormalities in bone formation or bone loss, conditions
XX CC that involve altered bone metabolism (e.g., idiopathic juvenile
XX CC osteoporosis), skeletal disease linked to breast cancer, mastocytosis,
XX CC Paget's syndrome of fibrous dysplasia, the present sequence is that of an
XX CC osteoblast differentiation associated cDNA marker of the invention.
XX CC Note: The sequence data for this patent did not form part of the printed
XX CC specification, but was obtained in electronic format directly from WIPO
XX CC at ftp.wipo.int/pub/published_pat_sequences.

XX PS Claim 1; SEQ ID NO 89; 78pp + Sequence Listing; English.

XX CC The invention relates to genes and their expression profiles are used
XX CC for:

XX CC (a) screening modulators of precursor stem cell differentiation into
XX CC osteoblasts, or bone tissue deposition;
XX CC (b) diagnosing abnormal deposition of bone tissue, abnormal rate of
XX CC osteoblast formation or osteoporosis; or
XX CC (c) treating or monitoring treatment of the conditions cited in (b), or
XX CC monitoring the progression of bone tissue deposition.
XX CC Specific conditions include postmenopausal osteoporosis, glucocorticoid
XX CC osteoporosis or male osteoporosis, osteopenia, osteodystrophy,
XX CC drug-induced abnormalities in bone formation or bone loss, conditions
XX CC that involve altered bone metabolism (e.g., idiopathic juvenile
XX CC osteoporosis), skeletal disease linked to breast cancer, mastocytosis,
XX CC Paget's syndrome of fibrous dysplasia, the present sequence is that of an
XX CC osteoblast differentiation associated cDNA marker of the invention.
XX CC Note: The sequence data for this patent did not form part of the printed
XX CC specification, but was obtained in electronic format directly from WIPO
XX CC at ftp.wipo.int/pub/published_pat_sequences.

XX SQ Sequence 3072 BP; 846 A; 688 C; 868 G; 670 T; 0 other;

Alignment Scores:
Pred. No.: 0.0057 Length: 3072
Score: 65.00 Matches: 13
Percent Similarity: 100.00% Conservative: 0

Percent Similarity: 100.00%
Best Local Similarity: 100.00%
Query Match: 24
Indels: 0
Gaps: 0

Score: 65.00
Percent Similarity: 100.00%
Best Local Similarity: 100.00%
Query Match: 21
Indels: 0
Gaps: 0

US-09-856-070-19 (1-13) x AAK98182 (1-3072)

QY 1 LysClnClnLeuMetLeuArqLeuGlnAspTyrGluCln 13
|||||
DB 1163 AAGACAGACTGATGCTGAGGCTGACAGGACTATGACAGAG 1201

RESULT 9

ID AAK98113
ID AAK98113 standard; cDNA: 3115 BP.

XX AAK98113;

XX 09-MAR-2001 (first entry)

XX Human colon cancer antigen nucleotide sequence SEQ ID NO:123.
XX Human; colon cancer; antigen; diagnosis; detection;
XX identification; cytostatic; cardioactive; neuroprotective; vulnary;
XX immunomodulatory; muscular; anaecological; gastrointestinal;
XX nephrotropic; antinecrotic; antibacterial; gene therapy; wound;
XX neural disorder; immune system disorder; muscular disorder;
XX reproductive disorder; gastrointestinal disorder; renal disorder;
XX infectious disease; cardiovascular disorder; ss.

XX Homo sapiens.

XX W020055451 A1.

XX 21-SEP-2000.

XX 08-MAR-2000; 2000W-US05883

XX 12-MAR-1999; 9905-0124270.

XX (HUMA-) HUMAN GENOME SCI INC.

XX Rosen CA, Ruben SM;

XX WPI: 2000 587544/55

XX P-IPDH; AAB53356.

XX Colon cancer associated gene sequences, referred to as colon cancer
XX antigens, useful for the treatment, prevention, and diagnosis of colon
XX disorders such as colon cancer -
XX Claim 1: Page 554-560; 2104pp; English
XX AAK97991 to AAK98763 encode the human colon cancer associated proteins
XX called human colon cancer antigens, given in AAK5323 to AAK5406. The
XX human colon cancer antigen can have cytostatic, cardioactive, muscular;
XX neuroprotective immunomodulatory anaecological, gastrointestinal,
XX vulnary, nephrotropic, antinecrotic and antibacterial activities, and
XX can be used in gene therapy. The colon cancer antigen polynucleotides,
XX proteins and antibodies to the proteins are useful for the prevention,
XX treatment and diagnosis of colon disorders, such as colon cancer. The
XX polynucleotides may be used in diagnostics and research, such as for
XX chromosome identification, and as hybridisation probes. The proteins
XX may also be used to prevent diseases such as neural disorders, immu-
XX system disorders, muscular disorders, reproductive disorders,
XX gastrointestinal disorders, wounds, renal disorders, infectious
XX diseases, and cardiovascular disorders. AAK98764 to AAK98772 and
XX AAK54007 represent sequences used in the exemplification of the present
XX invention.

XX Sequence 3115, 407, 873 A; 666 C; 872 G; 670 T; 4 other;

XX Alignment Scores:

XX Pred. No.: 0.00585

XX Length: 3115

US-09-856-070-19 (1-13) x AAK98113 (1-3115)
QY 1 LysClnClnClnMetLeuArqLeuGlnAspTyrGluCln 13
|||||
DB 1179 AAGACAGACTGATGCTGAGGCTGACAGGACTATGACAG 1217

RESULT 10

AAK70537/C

ID AAK70537 standard; DNA: 11445 BP.

XX AAK70537;

XX 06-NOV-2001 (first entry)

XX Human immune/haematopoietic antigen genomic sequence SEQ ID NO:25349.
XX Human; immune; haematopoietic; immune/haematopoietic antigen; cancer;
XX cytostatic; gene therapy; vaccine; metastasis; ds.
XX Homo sapiens.
XX W0200157182 A2.

XX 09-AUG-2001.

XX 17-JAN-2001; 2001W-US01354.

XX 31-JAN-2000; 2000US-0174065.

XX 04-FEB-2000; 2000US-0180628.

XX 24-FEB-2000; 2000US-0184664.

XX 02-MAR-2000; 2000US-0186350.

XX 16-MAR-2000; 2000US-0189874.

XX 17-MAR-2000; 2000US-0190076.

XX 18-APR-2000; 2000US-0198123.

XX 19-MAY-2000; 2000US-0205515.

XX 07-JUN-2000; 2000US-0209467.

XX 28-JUN-2000; 2000US-0214886.

XX 30-JUN-2000; 2000US-0215135.

XX 07-JUL-2000; 2000US-0216647.

XX 07-JUL-2000; 2000US-0216880.

XX 11-JUL-2000; 2000US-0217487.

XX 11-JUL-2000; 2000US-0217496.

XX 14-JUL-2000; 2000US-0218290.

XX 26-JUL-2000; 2000US-0220963.

XX 26-JUL-2000; 2000US-0220964.

XX 14-AUG-2000; 2000US-0224518.

XX 14-AUG-2000; 2000US-0224519.

XX 14-AUG-2000; 2000US-0225213.

XX 14-AUG-2000; 2000US-0225214.

XX 14-AUG-2000; 2000US-0225266.

XX 14-AUG-2000; 2000US-0225267.

XX 14-AUG-2000; 2000US-0225268.

XX 14-AUG-2000; 2000US-0225270.

XX 14-AUG-2000; 2000US-0225447.

XX 14-AUG-2000; 2000US-0225757.

XX 14-AUG-2000; 2000US-0225758.

XX 14-AUG-2000; 2000US-0225759.

XX 18-AUG-2000; 2000US-0226279.

XX 22-AUG-2000; 2000US-0226681.

XX 22-AUG-2000; 2000US-0226868.

XX 22-AUG-2000; 2000US-0227182.

XX 23-AUG-2000; 2000US-0227009.

XX 30-AUG-2000; 2000US-0228924.

XX 01-SEP-2000; 2000US-0229287.

XX 01-SEP-2000; 2000US-0229343.

XX 01-SEP-2000; 2000US-0229444.

XX 01-SEP-2000; 2000US-0229445.

RESULT 13

ABA74089/c

ID ABA74089 standard; DNA: 224 BP.

XX
AC ABA74089;XX
DI 01-FEB-2002 (first entry)XX
DE Human foetal liver single exon nucleic acid probe #22304XX
KW Human; foetal liver; gene expression, single exon nucleic acid probe; ss.XX
OS Homo sapiens.XX
PN WO200157277-A2.XX
PO 09-AUG-2001.XX
PP 30-JAN-2001; 2001WO-US00669XX
PR 04-FEB-2000; 2000US-0140312XX
PR 26-MAY-2000; 2000US-0207456.XX
PR 30-JUN-2000; 2000US-0608408.XX
PR 03-AUG-2000; 2000US-0142466.XX
PR 21-SEP-2000; 2000US-0234687.XX
PR 27-SEP-2000; 2000US-0236359.XX
PR 04-OCT-2000; 2000GB-0024263XX
PA (MOLIF-) MOLECULAR DYNAMICS INC.XX
PI Penn SG, Hanzel DK, Chen W, Rank DR;XX
XX WPI: 2001-483447/52.XX
PT Human genome-derived single exon nucleic acid probes useful for
XX analyzing gene expression in human foetal liver.XX
PS Claim 4; SEQ ID NO 22304; 60pp + sequence listing, English.XX
CC The invention relates to a single exon nucleic acid probe for
CC measuring human gene expression in a sample derived from human foetal
CC liver. The single exon nucleic acid probes may be used for predicting,
CC measuring and displaying gene expression in samples derived from human
CC foetal liver. The present sequence is a single exon nucleic acid
CC probe of the invention.XX
CC Note: The sequence data for this patent did not form part of the
CC printed specification, but was obtained in electronic format directly
CC from WIPO at ftp.wipo.int/pub/published_pat_sequences.XX
SQ Sequence 224 BP; 58 A; 63 C; 45 G; 58 T; 0 other.

Alignment Scores:

Pred. No.:	24.6	Length:	224
Score:	40.00	Matches:	7
Percent Similarity:	84.62%	Conservative:	4
Best Local Similarity:	53.85%	Mismatches:	2
Query Match:	61.54%	Indels:	0
DB:	22	Gaps:	0

US-09-856-070-19 (1-13) x ABA74089 (1-224)

QY 1 LysGluGluLeuMetLeuArgLeuGlnAspTyrGluGlu 13

||||| ||||| ||||| ||||| |||||

DB 57 AAGAGAAATCTGTTCTGGAAATCTGGTAAATATATAGGAAA 19

RESULT 14

AAK22545/c

ID AAK22545 standard; DNA: 224 BP.

XX
AC AAK22545;XX
DI 05-NOV-2001 (first entry)

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DE

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KW

KW

KW

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OS

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PN

PO

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PP

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PR

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PR

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PA

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SQ

Sequence 224 BP; 58 A, 63 C, 45 G, 58 T, 0 Other,

Alignment Scores:

Pred. No.:	24.6	Length:	224
Score:	40.00	Matches:	7
Percent Similarity:	84.62%	Conservative:	4
Best Local Similarity:	53.85%	Mismatches:	2
Query Match:	61.54%	Indels:	0
DB:	22	Gaps:	0

US-09-856-070-19 (1-13) x AAK22545 (1-224)

QY 1 LysGluGluLeuMetLeuArgLeuGlnAspTyrGluGlu 13

||||| ||||| ||||| ||||| |||||

DB 57 AAGAGAAATCTGTTCTGGAAATCTGGTAAATATATAGGAAA 19

RESULT 15

AAH61592/c

ID ABA61592 standard; DNA, 520 BP.

XX
AC ABA61592;XX
DI 01-FEB-2002 (first entry)XX
DE Human foetal liver single exon nucleic acid probe #9897.XX
KW Human; foetal liver; gene expression, single exon nucleic acid probe; ss.XX
OS Homo sapiens.XX
PN WO200157277-A2.

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XX 09 AUG-2001.
XX 10 JAN-2001: 2001WO-US000569.
XX 04 FEB-2000: 2000US-0180312.
XX 26 MAY-2000: 2000US-0207456.
XX 30 JUN-2000: 2000US-0608408.
XX 03 AUG-2000: 2000US-0632366.
XX 21 SEP-2000: 2000US-0244687.
XX 27 SEP-2000: 2000US-0236359.
XX 04 OCT-2000: 2000US-0024263.
XX (MOLE) MOLECULAR DYNAMICS INC.
XX Penn St, Hanzel DK, Chen W, Rank DR;
XX WPI; 2001-48447/52.
XX Human genome-derived single exon nucleic acid probes useful for
XX analyzing gene expression in human fetal liver.
XX Claim 1: SEQ ID NO 9897; 639pp + sequence listing; English.
XX the invention relates to a single exon nucleic acid probe for
XX measuring human gene expression in a sample derived from human fetal
XX liver. The single exon nucleic acid probes may be used for predicting,
XX measuring and displaying gene expression in samples derived from human
XX fetal liver. The present sequence is a single exon nucleic acid
XX probe of the invention.
XX Note: The sequence data for this patent did not form part of the
XX printed specification, but was obtained in electronic format directly
XX from WIPO at ftp.wipo.int/pub/published_pat_sequences.
XX Sequence 520 BP; 135 A; 145 C; 88 G; 152 T; 0 other;

Alignment Scores:
Pred. No.: 66.1 Length: 520
Score: 40.00 Matches: 7
Percent Similarity: 84.62% Conservative: 4
Best Local Similarity: 53.85% Mismatches: 2
Query Match: 61.54% Indels: 0
DB: 22 Gaps: 0

US 09 856-070 19 (1-13) x ABA61592 (1-520)
QY 1 LysGluLeuLeuMetLeuArgLeuGluAspTyrGluGlu 13
DB 150 AAGVAGAAATCTGTTCCTCGAATCTCCGTAACATGAGGAA 112

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Search completed: January 16, 2003, 17:19:45
Job time : 218.954 secs